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**Insensitivity of soil biological communities to phosphorus fertilisation in
intensively-managed grassland systems**

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Abstract

Efficient soil P cycling is essential for promoting optimal and sustainable grassland
growth. The soil biological community is regarded as an important source of available P
to the plant community. However, the effects of P fertilisation on the soil biota are
unclear. This study aimed to investigate the effects of P fertilisation on plant and soil

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biological communities in two intensively managed grassland sites that had been receiving mineral-based P fertilisation over a 14-year period. Both pastures had been frequently cut and harvested for plant material and only received inorganic fertilisers since the establishment of the trial. Both sites were sampled four times from October 2009 to April 2011 and plant and a range of soil biological parameters were determined at each sampling period. The main findings of this study showed that soil chemical measures, such as labile inorganic P and total P concentrations, and plant yield and P contents responded as expected to P fertilisation. However, all soil biological parameters either showed no response or inconsistent responses to P fertilisation over the experimental period. This study indicates that intensive management regimes, for example intensive plant harvesting and fertiliser regimes, appear to over-ride the relationship between plant and soil biological communities with respect to their response to P fertilisation, and thus their productivity is apparently not predicated upon biotic activity.

Keywords

Phosphorus – Phosphorus fertilisation – Microbial biomass phosphorus – Plant phosphorus contents – Harvesting regime – Grassland systems

Introduction

The application of fertilisers to grassland systems is a common management practise that aims to increase growth and productivity of the vegetation. There are different fertiliser types that function by supplying specific nutrients to the soil. One such type includes inorganic P-based fertilisers that are commonly applied to adjust the P status of the grassland soil (Bunemann et al., 2011). When applied, such fertilisers produce a flush of bioavailable P within a short timeframe in the immediate vicinity of the area of application (Stewart, 1991; Stevenson and Cole, 1999). This flush of bioavailable P disrupts the equilibrium between P in the mobile phase (water-soluble bioavailable P) and the solid phase (bound P), which ultimately develops into the adsorption of water-soluble P to components of the soil fabric, rendering it unavailable for uptake by plants and soil microorganisms. This inability of the plant community to readily assimilate P that has become bound to the soil matrix results in either the impairment of grass growth or the over-compensation of applied fertiliser P by the farmer to ensure that enough bioavailable P is supplied to the vegetation to optimise such growth (Stewart, 1991).

Whilst soil chemical processes can affect P availability, the soil biological community is also regarded as an important factor governing the availability of P to plants (Brookes, 2001; Achat et al., 2010). For example, the turnover of microbial biomass has been shown to release labile pools of organic P which can be mineralised and ultimately utilised by grassland swards. Based on this manner of P cycling, the biomass can be viewed as a potential slow release pool of labile organic P into the soil that can be accessed by plants (Brookes, 2001). Nevertheless, the actual size and activity of the

microbial biomass is dependent on multiple soil factors, such as the presence of other soil organisms, moisture content and soil type and the availability of other soil nutrients (Gregorich et al., 1991; Thomsen et al., 2003; Demoling et al., 2007; Chapin III et al., 2011). The turnover of the microbial biomass is critically linked to the activity of other soil organisms. For example, soil nematode communities exhibit a range of trophic interactions with other soil organisms. Some species feed exclusively on bacteria, some on fungi and others are plant parasites (Ingham et al., 1985). By feeding on microbial communities, nematodes have been shown to stimulate growth in the microbial community and facilitate the release of soil nutrients, particularly N, from the microbial biomass into the soil matrix (Ingham et al., 1985; Griffiths, 1994). By grazing on the microbial community, the turnover rate of microbial biomass increases and released nutrients aid in the synthesis of both microbial and plant biomass.

As stated above, soil type affects soil microbial communities. For example, increasing clay content has been reported to increase the biomass nutrient pool size of the microbial community due to a reduction in biomass turnover and an increase in soil moisture content (Gregorich et al., 1991; Thomsen et al., 2003). A reduction in biomass turnover occurs since increasing clay content facilitates the sorption of organic compounds to the soil matrix, thereby increasing the resistance of the soil organic matter pool to decomposition. Therefore, with increasing moisture content and a more decomposition-resistant organic matter pool, the soil microbial community is less active and biomass turnover occurs more slowly, which can result in the determination of greater microbial biomass sizes with increasing clay contents. Like the effects of soil type on biomass size, the size of the microbial biomass is also limited by the availability

of soil nutrients. One of the main nutrients known to be limiting microbial growth is C, which in turn limits assimilation of P (and other elements) (Demoling et al., 2007; Bunemann et al., 2011). With respect to C input into the soil, plants are recognised as being the primary source of C to the soil microbial community (Bardgett, 2005). Through rhizodeposition and plant senescence, plants supply different forms of C both directly to the soil surface via litter deposition (which may then be incorporated via the action of biota such as earthworms) and within the soil matrix via rhizodeposition. Such input of C stimulates the soil heterotrophic community and aids in soil nutrient cycling.

Whilst there are clear effects of soil biological communities on soil P cycling in the literature, the effects of P fertiliser application on soil biological communities are less apparent (Yeates, 1976; Ross et al., 1995; Sarathchandra et al., 2001; Gichangi et al., 2009). This ambiguity arises since varied results of P fertilisation have been reported on microbial and nematode communities. One common theme observed between these studies is linked to the joint application of both organic and inorganic fertiliser (Gichangi et al., 2009). The application of both forms has been shown to increase microbial biomass P concentrations above those observed from a sole application of inorganic fertiliser. However, when studies have investigated the sole application of inorganic P fertiliser, they either report inconsistent effects of P fertiliser on microbial biomass P over time or no effects at all (Ross et al., 1995; Sarathchandra et al., 2001).

With respect to nematode communities, responses to P fertilisation appeared to be coupled to livestock grazing, arising via greater nutrient input into the soil as faecal material. This return of organic nutrients to the soil stimulates soil microbial growth, which in turn increases nematode numbers (Yeates, 1976).

With equivocal data from the literature regarding P fertilisation effects on soil biology under pastures, a study was conducted which utilised different soil types that had received long-term applications of inorganic P fertiliser (>14 years). The aim of this study was to investigate and to further clarify how plant and soil biological communities were affected by inorganic P fertiliser applications. We hypothesised that both plant and soil communities will respond to P fertilisation, and specifically that the application of P fertiliser will promote greater plant P yields and microbial biomass P concentrations.

Methods

Field description and sampling

All sampling was carried out on two long term grassland sites located in the dairy farm area at Johnstown Castle, Wexford, County Wexford, Ireland (latitude 52 °17 N, longitude 06°30 W) (Daly, 2005). Site 1 is founded on a fine loamy soil (22% coarse sand, 27% fine sand, 29% silt and 22% clay) and Site 2 on a coarse loamy soil (29% coarse sand, 22% fine sand, 33% silt and 16% clay). Both of these sites received four different P application rates, viz. 0, 15, 30 and 45 kg P ha⁻¹ y⁻¹ of 16% single superphosphate (2 CaSO₄ + Ca(H₂PO₄)₂) (hereafter denoted P0, P15, P30 and P45, respectively). P was applied to each plot in February each year. There were four replicates of each treatment within a site and, therefore, 16 plots per site. Each plot had the dimension 10 x 2 m². Different P application rates were randomly positioned within a site according to a randomised block design. The plant species present in both grassland sites were *Lolium perenne* L., *Poa trivialis* L., *Agropyron repens* (L.) Beauv., *Trifolium repens* L. and various *Agrostis* spp. Aboveground plant material in both sites

was cut eight times per year to a height of 5 to 6 cm using a plot harvester. After harvesting plant material in both sites, all plots received 40 kg N ha⁻¹, as calcium ammonium nitrate (5 Ca(NO₃)₂ NH₄NO₃). Potassium was also applied as muriate of potash (KCl) at a rate of 125 kg K ha⁻¹ y⁻¹ to compensate for potassium removal from both systems. These trials were set up in February 1995 and thus had been established for 14 years prior to the onset of this study.

Rainfall, air temperature and soil temperature were monitored over the experimental period of this study. Rainfall was determined using a tipping bucket rain gauge. Air temperature was measured using a dry platinum resistance thermometer. Soil temperature was measured at a depth of 10 cm using a platinum resistance thermometer.

Soil samples were collected from both sites in October 2009, April 2010, October 2010 and April 2011. Five sample cores (diameter 6 cm, depth 10 cm) were collected from each individual plot in a W-best-of-fit sampling design to ensure representative non-biased sampling and composited on a per-plot basis.. Samples were firstly broken by hand, homogenised and 100 g fresh weight subsampled for nematode analysis.

Remaining soil was coned, quartered and subsampled for 4 mm and 2 mm sieving to remove the majority of plant and stone material. Soil sieved to 4 mm was subsampled for phospholipid fatty acid analysis (PLFA) analysis and the soil that was sieved to 2 mm was subsampled for microbial biomass C, N and P. During the course of these experiments, all fresh soil samples for microbial biomass analyses were stored at 4°C. Samples for PLFA analysis were stored at -80°C using an ultra low temperature freezer and freeze dried. Different sieve sizes were used in this experiment since the aim was not to relate such properties but to examine how they were affected by P fertilisation.

By sieving to 4 mm for PLFA analysis, less disturbance of the soil matrix would occur compared to 2 mm sieving (Petersen and Klug, 1994). Sieving to 2 mm was chosen for all other parameters because this would enable for more comprehensive comparisons with the literature, especially in the case of microbial biomass C, N and P concentrations, since sieving to 2 mm is a more common practice for determining such concentrations (Sarathchandra et al., 2001; Turner et al., 2001; Lui et al., 2010; Lui et al., 2012). All remaining soil was dried at 40°C, sieved to 2 mm and used to determine soil pH, loss-on-ignition, Morgan extractable P and total P.

Plant dry matter yield and phosphorus content

All plant material harvested from each plot was dried at 70°C for 72 h and weighed to determine plant dry matter yield. Plant P contents were determined by digesting dried plant subsamples using a Gerhardt Kjeldatherm block digester and measured using a continuous flow peak height analyser (McCormack, 2002). Briefly, one No. 80 'Kjeltabs' tablet (Carl Stuart Limited, Republic of Ireland) was added to the dried plant sample, 5 ml of 95% sulphuric acid (H₂SO₄) and 3 ml of hydrogen peroxide (H₂O₂) were then added to the sample, heated at 150°C for 1 h and then at 390°C for 1.5 h using the block digester. When digested, samples were diluted with deionised water to make the sample volume up to 50 ml. Samples were then filtered and an aliquot was prepared for P measurement.

Plant P content was determined based on orthophosphate concentrations produced during digestion. Orthophosphate concentrations were determined by reacting

orthophosphate with a molybdovanadate reagent, which were then quantified
colourimetrically by reading absorbance at 420 nm (McCormack, 2002).

Soil chemical methodologies

Labile inorganic P was determined using the Morgan's extraction method (McCormack, 2002). Briefly, labile inorganic P was extracted from 3 g dry soil using 15 ml of a 0.62 M sodium hydroxide (NaOH) and 1.25 M acetic acid (CH₃COOH) solution adjusted to pH 4.8 at a 1:5 (w/v) soil to solution ratio (Peech and English, 1944; McCormack, 2002). Soils were shaken for 30 minutes on gyratory shaker and filtered. P concentrations were measured spectrophotometrically at 880 nm using the ammonium molybdate-ascorbic acid method (Murphy and Riley, 1962). Total P concentrations were determined in soil collected in April 2011 using the aqua regia digestion method (ISO 11466:1995). Soil samples were ball milled to less than <0.1 mm and 2 g of this milled material were weighed into digestion tubes. Both 16 ml of hydrochloric acid and 4 ml of nitric acid were added to each digestion tube, these tubes were then placed into a digestion block. Samples were digested at 140°C for 2 h, thereafter samples were filtered using No. 2 Whatman filter paper and the filtered extract made up to a final volume of 100 ml using 2 M nitric acid. Extracted P concentrations were measured using inductively-coupled plasma optical-emission spectroscopy. Soil pH was determined using a soil deionised water ratio of 1:2 (w/v) (McCormack, 2002). Organic matter contents were determined by loss-on-ignition (Rowell, 1994)

Soil biological methodologies

Prior to the determination of microbial biomass C, N and P, all soil was incubated at 21°C and field moisture content for 7 days. After this incubation period, microbial biomass C and N concentrations were determined following a modified fumigation extraction methodology (Brookes et al. 1985; Vance et al. 1987). Briefly, soil samples were weighed out in duplicate; one sample for fumigation and another for baseline C and N concentrations. Samples to be fumigated were placed into a vacuum desiccator with ethanol-free chloroform and fumigated for 24 h. Following fumigation, samples were extracted using 0.5 M potassium sulphate (K_2SO_4) (1:4 soil solution ratio) on a side-to-side shaker for 30 min. The other subsamples were not fumigated but were extracted using 0.5 M K_2SO_4 under the same conditions as the fumigated samples. After extraction, C and N concentrations were measured using a Shimadzu TOC-V and TNM-1 analyser (Shimadzu Corporation, Japan). Reported concentrations were corrected using the conversion factors stated in the literature. Microbial biomass C and N concentrations were estimated using the conversion factors 0.45 and 0.54, respectively (Brookes et al. 1985; Vance et al. 1987).

Microbial biomass P concentrations were determined following the fumigation extraction procedure (Brookes et al. 1982). Samples were weighed in triplicate; one for fumigation, another for baseline P concentrations and a third to compensate for P adsorption during the extraction process. The fumigation sample was fumigated using ethanol-free chloroform in a vacuum desiccator for 24 h. Following fumigation, P was extracted from fumigated and non-fumigated subsamples using a 0.5 M sodium bicarbonate ($NaHCO_3$) solution adjusted to pH 8.5 at a 1:20 soil solution ratio. The third sample, to account for P adsorption, was also extracted using 0.5 M $NaHCO_3$ but an

additional 1 ml of 25 $\mu\text{g P g}^{-1}$ was added to the sample: 1 ml of deionised water was added to fumigated and non-fumigated samples. P was applied as potassium dihydrogen orthophosphate (KH_2PO_4). All samples were shaken for 30 min following the addition of the extractant on a side-to-side shaker. Once extracted, P concentrations were measured by UV spectroscopy at 880 nm following the ammonium molybdate-ascorbic acid method (Murphy and Riley, 1962). Microbial biomass P was calculated using the equation stated in Brookes et al. (1982).

PLFA analysis was determined using a modified procedure described by Frostegard et al. (1991), which was based on the method described by Bligh and Dyer (1959). Lipids were extracted from freeze dried samples using a mono-phase Bligh and Dyer extractant. This reagent consisted of chloroform, methanol and 0.15 M citrate buffer (1:2:0.8 v/v/v, respectively). With this extractant, samples were sonicated for 30 min and then shaken for a further 30 min before being centrifuged. Using the organic top layer of the extractant, phospholipids were separated from neutral lipids and glycolipids in SPE cartridges (Silica Cartridge Sep Pak Vac 3 cc (500 mg), Waters Scientific, USA) by solid phase extraction. Phospholipids were subjected to mild alkaline methanolysis to form fatty acid methyl esters, which were then separated using gas chromatography and detected using a flame ionisation detector set to 310°C (Varian, USA). The column used was a HP-5 (Agilent Technologies, USA) capillary column. A 26 Bacterial Acid Methyl Ester mix was used as an external standard to identify commonly occurring PLFA markers in the sample.

Nematodes were washed from 100 g fresh weight of soil using an Oostenbrink elutriator (MEKU Erich Pollähne GmbH, Germany) (Seinhorst, 1962). This form of elutriation

passes a sample suspended in water through a series of sieves (180, 120, 95 and 52 μm apertures). Nematodes were separated from elutriated material using Baermann funnels over a 48 hr period. Following this period, 50 ml of sample was collected from the funnel. Samples were left for one day, after which 45 ml was removed. Constant checks were made to ensure that no nematodes were removed during this process. The remaining 5 ml of sample was mixed until nematodes were re-suspended and 2 ml transferred to a counting dish, to generate a representative measure of the total number of nematodes in the sample. The number of nematodes present in this 2 ml subsample was then counted using a light microscope. The abundance of nematodes (number g^{-1} dry soil) within the sample was then estimated based upon this representative subsample.

Earthworms were extracted using the Octet method (Schmidt, 2001) in April and October 2010. Vegetation was clipped to the soil surface and eight stainless steel electrodes were pushed into the soil (40 cm deep) at marked positions around a sampling ring (area 0.125 m^2). The stainless steel rods were joined to the sampling ring using the designated connector clips and the octet machine attached to a 105 AHR 12-V car battery. The machine was turned on and voltage increased from 200/250 V to 500/600 V at 5 min intervals for the first four steps and 10 min for the remaining two steps. Earthworms were collected from within the ring when they fully emerged. Live earthworms were then weighed, inclusive of gut content.

Statistics

Data were analysed using factorial repeated measures ANOVA via Statistica v.9 (Stats Soft, 2010). Site, P fertilisation rate and sampling occasion were designated as main effects, with sampling occasion as the repeated measure level. Total P data was analysed using factorial two-way ANOVA since data was only obtained in April 2011 and not a repeated measure. With respect to microbial community structure, all PLFA markers were normalised to mol% within each sample and analysed using principal component analysis (PCA) to produce principal components (PC) for the whole data-set. Resultant PCs were subsequently analysed using factorial repeated measures ANOVA to investigate treatment effects as stated above.

Results

Environmental conditions

With respect to the sampling occasions adopted for this experiment, the soil and air temperatures were similar at all four occasions (Fig. 1 (a)). The range of soil temperatures observed for these occasions were between 12.8 ± 0.17 and $10.3^\circ\text{C} \pm 0.31$ S.E. Air temperature during these occasions ranged from 12.1 ± 0.26 to $8.5^\circ\text{C} \pm 0.36$ S.E. The greatest difference between soil and air temperatures occurred in April 2010 (soil temperature; 10.3 ± 0.31 , air temperature; $8.5^\circ\text{C} \pm 0.36$ S.E., whereas soil and air temperatures in October 2009, October 2010 and April 2011 were similar. Unlike soil and air temperatures, greater rainfall was observed in October 2009 and 2010 (5.2 ± 1.8 and $3 \text{ mm} \pm 0.8$ S.E., respectively) compared to April 2010 and 2011 ($0.9 \text{ mm} \pm 0.5$ S.E. for both dates) (Fig. 1 (b)).

Site and seasonal effects on soil chemical and biological parameters

For most measured chemical and biological parameters, greater quantities and concentrations were observed in the fine loamy soil compared to the coarse loamy soil. No interactions between P fertilisation and Site were observed on both plant P and plant dry matter contents ($p > 0.05$ for both parameters). When analysing plant P contents, a significant site effect was apparent ($p < 0.01$), which showed that greater plant P contents were manifest on the fine loamy compared to the coarse loamy soil (Table 1). No site effects were observed on plant dry matter contents (Table 1).

No interaction between P fertilisation and Site was apparent on total soil P concentrations ($p > 0.05$). However, a Site effect was observed ($p < 0.001$), which showed greater total P concentrations in the fine loamy compared to the coarse loamy soil (Fine loamy soil = 670; Coarse loamy soil = 517 mg kg⁻¹ ± 14.2 pooled S.E.). Morgan's P concentrations and organic matter contents were significantly affected by Site whereas pH was not (Table 2). However, these site effects on Morgan's P and organic matter differ between seasons, which were apparent by interactions with sampling occasion (Table 2). The interactions for both of these parameters showed greater concentrations and contents were measured in April 2010 and 2011 compared to October 2009 and 2010 (Fig. 2 (a) and Fig. 3). Whilst no Site effect was apparent for soil pH, a significant Site by Time interaction was observed (Table 2). This interaction indicated a general decrease in pH in both sites, but the decrease in pH started on different occasions for both sites. In the fine loamy soil, a consistent pH value was

324 observed until April 2010, thereafter pH decreased in October 2010 and decreased
325 further in April 2011 (October 2009 = 6.5 ± 0.09 ; April 2010 = 6.6 ± 0.07 ; October
326 2010 = 6.3 ± 0.07 ; April 2011 = 6.01 ± 0.06 S.E.). In the coarse loamy soil, a consistent
327 pH value was apparent until October 2010 after which pH decreased in April 2011
328 (October 2009 = 6.5 ± 0.09 ; April 2010 = 6.5 ± 0.07 ; October 2010 = 6.5 ± 0.07 ; April
329 2011 = 6.01 ± 0.06 S.E.).

330 Site effects were observed on microbial biomass C, N and P concentrations and on
331 microbial community structure (Table 3). In general, microbial biomass C, N and P
332 concentrations were greater in the fine loamy compared to coarse loamy soil. However,
333 such site effects on all microbial parameters were confounded by interactions with
334 sampling occasion. Microbial biomass C concentrations exhibited a complex third-order
335 interaction between Site, P fertilisation treatment and Time (Table 3 and Fig. 4).
336 Nevertheless, with respect to site temporal variation, there was greater variation in
337 microbial biomass C concentrations in the coarse loamy soil across the experimental
338 period compared to the fine loamy soil (Fig. 4). With respect to differences between
339 sampling occasions, both sites showed a relatively large increase in microbial biomass
340 C concentrations in April 2011 compared to all other sampling occasions. Microbial
341 biomass P concentrations increased in the fine loamy soil over the experimental period
342 (October 2009 = 47.4 ± 3.4 ; April 2010 = 70.7 ± 2.8 ; October 2010 = 66.9 ± 3.3 ; April
343 2011 = $90.4 \mu\text{g g}^{-1} \pm 2.9$ S.E.), whereas in the coarse loamy soil, an increase occurred
344 from October 2009 to April 2010, which was followed by consistent concentrations in
345 October 2010 and April 2011 that were similar to April 2010 (October 2009 = $34.4 \pm$
346 3.4 ; April 2010 = 50.4 ± 2.8 ; October 2010 = 50.5 ± 3.3 ; April 2011 = $52.4 \mu\text{g g}^{-1} \pm 2.9$

S.E.). With respect to microbial biomass N, greater concentrations were observed in the fine loamy soil in October 2009 and April 2011. In April 2010 and October 2010 similar concentrations were observed in both sites (Fine loamy soil: October 2009 = 184.7 ± 8.7 ; April 2010 = 100.6 ± 5.2 ; October 2010 = 152 ± 7.1 ; April 2011 = $212.3 \mu\text{g g}^{-1} \pm 4$ S.E. Coarse loamy soil: October 2009 = 123 ± 8.7 ; April 2010 = 93.8 ± 5.2 , October 2010 = 155 ± 7.1 ; April 2011 = $151 \mu\text{g g}^{-1} \pm 4$ S.E.). Microbial phenotypic community structure showed marked significant variation over the experimental period. Such differences were apparent in PC1-3 (Table 3 and Fig. 5).

The period in which soil was sampled also had significant effects on nematode abundance and earthworm biomass (Table 3). With respect to nematode abundance, a significant Site by Time interaction was observed. In the fine loamy soil, nematode abundance was greater in April 2010 and 2011 compared to October 2009 and 2010 (October 2009 = 33.7 ± 2.25 ; April 2010 = 50.3 ± 2.88 ; October 2010 = 33.6 ± 2.68 ; April 2011 = 50.3 individuals g^{-1} dry soil ± 2.88 S.E.). In the coarse loamy soil, nematode abundance was significantly reduced in October 2009 compared to all other sampling dates (October 2009 = 24.6 ± 2.25 ; April 2010 = 40.2 ± 2.88 ; October 2010 = 42.7 ± 2.68 ; April 2011 = 40.2 individuals g^{-1} dry soil ± 2.88 S.E.). Regarding earthworm biomass, greater biomass was revealed in October 2010 ($77.4 \text{ g m}^{-2} \pm 5.3$ S.E) compared to April 2010 ($55.3 \text{ g m}^{-2} \pm 4$ S.E). A significant Site effect was observed on earthworm biomass ($p < 0.01$), which showed greater biomass in the fine loamy soil ($77.3 \text{ g m}^{-2} \pm 4.9$ S.E.) compared to the coarse loamy soil ($55.4 \text{ g m}^{-2} \pm 4.9$ S.E.).

Phosphorus fertiliser effects on soil chemical and biological parameters

The application of P fertiliser at different rates to both sites had significant effects on plant dry matter and P contents ($p < 0.001$ for both parameters). With respect to dry matter content, the application of P fertiliser up to a rate of $30 \text{ kg P ha}^{-1} \text{ y}^{-1}$ significantly increased dry matter production. The application rate at $45 \text{ kg P ha}^{-1} \text{ y}^{-1}$ produced the same dry matter content as the $30 \text{ kg P ha}^{-1} \text{ y}^{-1}$ treatment. Plant P contents increased with increasing P fertilisation rate (Table 1).

Total soil P concentrations were significantly affected by P fertilisation rate ($p < 0.001$). The P fertilisation effect revealed that total P concentrations generally increased with increasing P fertilisation rate ($P_0 = 504$; $P_{15} = 566$; $P_{30} = 590$; $P_{45} = 714 \text{ mg kg}^{-1} \pm 20$ pooled S.E.). Specifically, *post hoc* analysis showed that total P concentrations in the P_0 treatment were significantly lower than all other treatments, whereas concentrations in the P_{45} treatment were significantly greater than all other treatments. No significant difference between the P_{15} and P_{30} treatments was apparent. Morgan's P concentrations showed a significant P fertilisation x sampling occasion interaction (Table 2), manifest as a general increase in Morgan's P with increasing P fertilisation rate. However, greater variability in P concentration across all sampling occasions was observed in the P_{45} treatment compared to all other P fertilisation regimes (Fig. 2 (b)). No P fertilisation effect was observed on soil organic matter content (Table 2).

No P fertiliser effect or interaction was observed on microbial biomass P and N concentrations, nematode abundance or earthworm biomass (Table 3). With respect to microbial biomass C, no consistent effects of P fertiliser were observed across both sites

and sampling occasions, as apparent by a third-order interaction (Table 3 and Fig. 4). This inconsistent effect revealed that different individual P treatments exhibited significantly greater or lower microbial biomass C concentrations at particular sampling occasions. Furthermore, microbial biomass C responses to P fertilisation across all sampling occasions differed between the two sites.

No consistent P fertilisation effects were observed on microbial phenotypic community structure. This inconsistent effect was manifest as a second-order interaction between P fertilisation and Time for PC1, 2 and 3 (Table 3). These interactions revealed that different individual P treatments were associated with community structures that were significantly different from all other P treatments at each sampling occasion (Fig. 5 (a) and (b)) No particular trends in microbial community structure are observed over time between the different P treatments.

Discussion

Seasonal and soil type effects on the soil biota in grassland systems

All biological properties varied significantly between seasons in this study, with microbial biomass C, N and P differing between all sampling periods, albeit with no consistent trends between them. Many other studies have similarly reported temporal variation in soil microbial properties (He et al., 1997; Krämer and Green, 2000; Chen et al., 2003). Seasonal variations in microbial nutrient pools has been associated with changes in soil moisture, soil temperature, root growth and activity (rhizodeposition) and organic matter input through plant senescence (Chen et al., 2003). In the study conducted by Chen et al. (2003), microbial biomass C and P were shown to vary

temporally, with greater variability observed in microbial biomass P than C. It appeared that microbial biomass P was more sensitive to plant growth (thus plant P demand) and soil moisture content compared to microbial biomass C. This result has also been observed in another study by Tate et al. (1991). In particular, moisture deficit in the soil may disrupt the diffusion of water-soluble P to the microbial community, thus affecting microbial assimilation of P on a temporal scale (He et al., 1997). He et al. (1997) explained that since a deficit in soil moisture content did not affect microbial biomass C, but coincided with a decrease in microbial biomass P, then this may represent a loss of P during microbial biomass P turnover and an inefficiency to then mobilise P from the soil. This effect of moisture deficiency on microbial biomass P may also be heightened by the activity of the plant community. The evapotranspiration induced by the plant community would not only hinder the diffusion of water-soluble P to the microbial community but could also be actively competing with the microbial community for smaller concentrations of orthophosphate in the soil solution (He et al., 1997). Therefore, the amount of P available to the microbial biomass may be severely limited at specific times of the year and may explain why microbial biomass P concentrations fluctuated during the experimental sampling period.

In this study, greater microbial biomass C, N and P was generally observed in the fine loamy soil compared to the coarse loamy soil. This finding complements previous studies focusing on textural effects upon C and N mineralisation and microbial biomass C and N (Gregorich et al., 1991; Hassink, 1994). Furthermore, an increase in soil organic matter with increasing clay content has also been reported (Gregorich et al., 1991). This complements soil organic matter behaviour in this study, since the fine

loamy soil contained both greater organic matter contents and more silt and clay compared to the coarse loamy soil (4% and 6% more silt and clay, respectively). Soil organic matter has been shown to be greater with increasing clay content due to the greater sorption of organic compounds to the clay components of the soil matrix (Gregorich et al., 1991). Such sorption would increase the stability of organic matter and retard soil decomposition processes. With reduced organic matter decomposition, the microbial biomass becomes less active which results in a slower biomass turnover rate. Therefore, greater microbial biomass C, N and P pools were observed in the fine loamy soil as this reflects a slower turnover and less active microbial community compared to the microbial biomass in the coarse loamy soil (Gregorich et al., 1991; Hassink, 1994).

Plant and microbial responses to phosphorus fertilisation

This study highlights that the application of P fertiliser affected plant P yields but not soil microbial biomass P concentrations, despite an increase in labile inorganic P (Morgan's P) with increasing P fertilisation. Therefore, the hypothesis of this study was rejected since no increase in microbial biomass P was observed despite an increase in plant P yields. Other studies assessing the effect of P fertilisation on soil microbial biomass have generally found mixed responses in these circumstances. Some studies focusing on the long-term application of inorganic fertiliser P to grassland systems have found no effects on microbial biomass (Sarathchandra et al., 1993; Sarathchandra et al., 2001), whilst other studies have reported mixed responses to P fertilisation. Ross et al. (1995) presented results that showed inconsistent effects of rock phosphate on microbial biomass P concentrations in a New Zealand pasture which was analysed over a two year

period. This study conducted by Ross et al. (1995) found that microbial biomass P did respond to P fertilisation on some sampling occasions, but not all. The application of inorganic P fertiliser has also been shown to decrease microbial biomass C (Parfitt et al., 2010). The explanation for this observation was attributed to a shift in microbial community structure, since a more bacterial dominated community occurred in the presence of a high fertility system that received inorganic P fertiliser. Consequently, a more bacterial dominated community contributed to a lower microbial biomass C concentration due to a more rapid turnover of such biomass due to grazing by micro-fauna. In this present study, no consistent P fertilisation effects were observed on the soil biota. Therefore, the absence of an effect in this instance does complement some results from other grassland systems that have focused on the sole application of inorganic P fertilisers.

In contrast, other studies have reported stimulatory effects of P fertilisation on the microbial biomass P concentrations. In a maize-wheat crop rotation system, Lui et al. (2010) reported that the application of inorganic P fertiliser combined with inorganic N fertiliser increased microbial biomass P concentrations. It was, however, highlighted that the greatest increase in microbial biomass occurred when inorganic and organic fertiliser were applied together. In another study conducted by Parfitt et al. (2005), P fertilisation effects were also observed on microbial biomass P concentrations in New Zealand pastures that were grazed by sheep under both conventional and organic fertiliser management. The P fertilisation effect was attributed to P limitation in the control treatment, whilst no differences were observed between fertiliser management types. One common theme that appears to be associated with the P fertiliser effects on

the soil biota is the importance of fertiliser type and management. When effects are observed, they were seemingly stimulated particularly by the input of organic material. In contrast, the fertiliser type and grassland management regime adopted in this present study did not involve the input of organic matter into the system, since only inorganic fertilisers were applied and no livestock were present. Furthermore, the intensive harvesting regime adopted in this study, involving frequent and efficient removal of plant biomass from the system, would have certainly curtailed plant-mediated inputs into the soil. With respect to the importance of plant litter inputs to stimulate P demand in the microbial community, a study by Liu et al. (2012) investigated how P additions to forest systems under different management regimes affected microbial biomass P. Liu et al. (2012) showed that the addition of P to a forest system that was not disturbed increased both microbial biomass C and P, whereas P addition to a disturbed forest system failed to invoke a response. It was concluded that C and N limitation in the disturbed forest scenario may have limited P acquisition due to an absence of nutrient accumulation. Therefore, we propose that one possible explanation for microbial biomass P not responding to P fertilisation in the present study may be linked to the management of these grassland sites, since such intensive harvesting may have governed the availability of other soil nutrients. The removal of aboveground biomass over a 16 year period would have constantly been removing C from the system and manipulated plant behaviour (Nevens and Rehuel, 2003; Vinther, 2006; Ilmarinen and Mikola, 2009; Ilmarinen et al., 2009). Cutting in this manner over such a long period of time may have promoted similar rhizodeposition patterns across all P fertilisation treatments since a greater investment in aboveground plant dry matter and P yields were

apparent with increasing P fertilisation. Therefore, by intensively cutting these grassland sites, the response of soil biological communities to P fertilisation may be lost as the plant community is the primary source of labile C to soil microbial communities (Bardgett, 2005). This impact of similar C inputs may be manifest in this study through the inconsistent effects of P fertilisation on microbial biomass C and microbial community structure. In this context, the interaction between plant and microbial communities in these grassland systems may have been driven by a similar input of plant material across all P fertilisation treatments, which would have limited the potential stimulatory effects of plant inputs on the microbial community in the presence of P fertiliser.

Despite intensive C harvesting from these sites, microbial biomass C concentrations were not low compared to other studies in grassland systems (Turner et al., 2001). Turner et al. (2001) looked at microbial biomass C concentrations in a range of soil types in 29 UK permanent grasslands. These concentrations ranged from 412 ± 19 to $3412 \pm 21 \mu\text{g g}^{-1}$. The largest concentrations observed were derived from soils with low percentage sand contents and high clay contents. Larger concentrations of microbial biomass C in the two grassland sites investigated in this study indicated that C may not be the limiting nutrient. Further evidence for this was supported by soil loss-on-ignition contents, which was similar to those observed in other Irish grasslands (Bourke et al., 2008). Bourke et al. (2008) reported mean loss-on-ignition soil contents of 8.22% (ranging from 6 - 10.9%) in sampled grassland sites at the Johnstown Castle Estate. Thus, the accumulation of organic material in these two grassland sites was not reduced by the cutting regime despite its intensity. This may support the conclusion that it was

the consistent inputs of similar quantities and qualities of C across all P fertilisation regimes that was limiting microbial P acquisition in plots that received P fertilisation, since no P fertilisation effect was observed on soil organic matter contents in this study.

Conclusions

From an agronomic perspective, both of these sites responded as expected to different rates of P fertilisation. In particular, both sites showed that labile and total soil P concentrations and plant yields and P contents increased with increasing P fertilisation. However, despite consistent effects on soil P concentrations and the plant community, no consistent P fertiliser effects were observed on soil biological communities. Such differences between plant and soil biological communities may be linked to the management of these grassland sites, since both sites were intensively managed via exclusively inorganic mineral nutrient inputs and extensive removal of plant biomass. This may have manipulated the interactions between plant and soil biological communities, which could have potentially limited the acquisition of P by the soil microbial community. One such management technique that may have greatly affected the interaction both communities is the adopted harvesting regime. By adopting an intense harvesting regime, the input of plant material into the soil may have been similar across all P fertiliser treatments, resulting in different responses between plant and soil biological communities. This study indicates that intensive management regimes appear to decouple the relationship between plant and soil biological communities with respect to their response to P fertilisation, and thus their productivity is apparently not predicated upon soil biotic activity. The extent to which grasslands are then able to carry any capacity to function when such fertiliser inputs cease is unclear, which is

critical considering the expense of P fertilisers to the agricultural community. Further work is still required to truly define the interactive effects of multiple grassland management techniques on the interactions between plant and soil biological communities. By fully understanding these interactions, it may be possible to promote greater P cycling in the soil, via the soil biota, which may open new insights into reducing plant dependency on fertiliser applications.

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692

Figure captions

Fig. 1 Environmental conditions in the Johnstown Castle estate from October 2009 to April 2011, (a) denotes air and soil temperature and (b) denotes rainfall. Error bars denote standard error.

Fig. 2 Mean Morgan's P concentrations expressed by (a) the interaction between sites and sampling periods and (b) the interaction between P fertilisation rates and sampling periods. Same letters denote homogeneous means ($p < 0.05$) using Fisher least significant difference test. Error bars denote standard error.

Fig. 3 Interactions between sites and sampling periods with respect to soil organic matter content. Same letters denote homogeneous means ($p < 0.05$) using Fisher least significant difference test. Error bars denote standard error.

Fig. 4 Interactions between site, sampling period and P fertilisation (Abbreviated as P: numerals denote application rate $\text{kg}^{-1} \text{ha}^{-1} \text{y}^{-1}$) on microbial biomass C concentrations (Third-order interaction significance $p = 0.028$). There are 15 homogenous mean groups associated with this interaction, hence letters denoting these groups have been omitted for clarity. Error bars denote pooled standard error.

Fig. 5 Differences in phenotypic microbial community structure in the presence of four different P fertilisation regimes (Abbreviated as P: numerals denote application rate $\text{kg}^{-1} \text{ha}^{-1} \text{y}^{-1}$) at four sampling periods as expressed by (a) PC1 and PC2 and (b) PC2 and PC3. Percentage variation accounted for by respective PCs shown in parenthesis. The bounding ellipses drawn around each sampling time are to assist visualisation, and have no formal statistical derivation.

Table 1 Mean total plant dry matter yields and total plant P contents over the course of this study in the presence of four P fertiliser regimes and across two grassland sites. Site 1 is a fine loamy soil and Site 2 is a coarse loamy soil. Same Letters denote homogeneous means ($p < 0.05$) using Fisher least significant difference test. Standard error is represented as error mean squares (EMS).

P fertilisation rate (kg P ha⁻¹ y⁻¹)	(kg m²)	(g m²)
	Total dry matter	Total plant P
0	1.38 a	1.87 a
15	1.67 b	2.93 b
30	1.79 c	3.86 c
45	1.79 c	4.42 d
EMS	0.01	0.07
Site		
1 (Fine loam)	1.67 a	3.44 a
2 (Sandy loam)	1.64 a	3.10 b
EMS	0.01	0.07

Table 2 ANOVA table showing the effects and interactions of Site, P fertilisation rate and sampling occasion (Time) on general soil parameters.

	Morgan's P ($\mu\text{g g}^{-1}$)	pH	Loss-On-Ignition (%)
Site	***	-	***
P treatment	***	-	-
Time	***	***	***
Site x Time	**	*	***
P treatment x Time	***	-	-
Site x P treatment	-	-	-
Site x P treatment x Time	*	-	-

Significance terms denote *** $p < 0.001$; ** $p < 0.01$; * $p < 0.05$; - $p > 0.05$, respectively.

728 **Table 3** ANOVA table showing the effects and interactions of Site, P fertilisation rate and sampling
 729 occasion (Time) on measured soil biological parameters.

	Microbial biomass ($\mu\text{g g}^{-1}$)			Microbial community structure			Soil faunal	
	C	N	P	PC1 (38%)	PC2 (16%)	PC3 (11%)	Earthworm biomass (g m^{-2})	Nematode abundance (number g^{-1} dry weight)
Site	***	***	***	-	***	***	**	-
P treatment	**	-	-	*	-	-	-	-
Time	***	***	***	***	***	***	**	***
Site x Time	***	***	***	-	-	-	-	***
P treatment x Time	**	-	-	*	**	*	-	-
Site x P treatment	-	-	-	-	-	-	-	-
Site x P treatment x Time	***	-	-	-	-	-	-	-

730

731 Significance terms denote *** $p < 0.001$; ** $p < 0.01$; * $p < 0.05$; - $p > 0.05$, respectively.

732

733

Figure 1

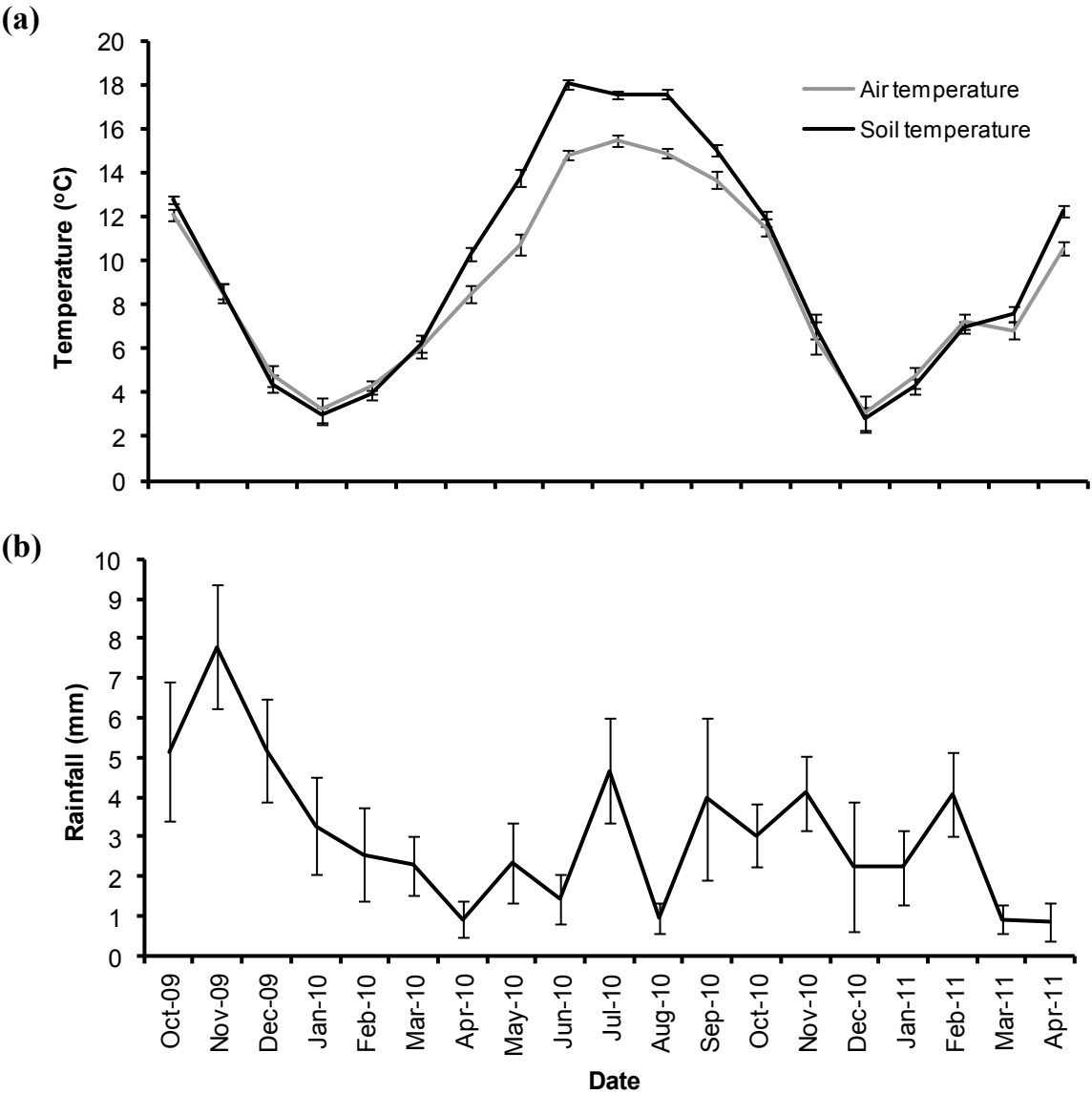
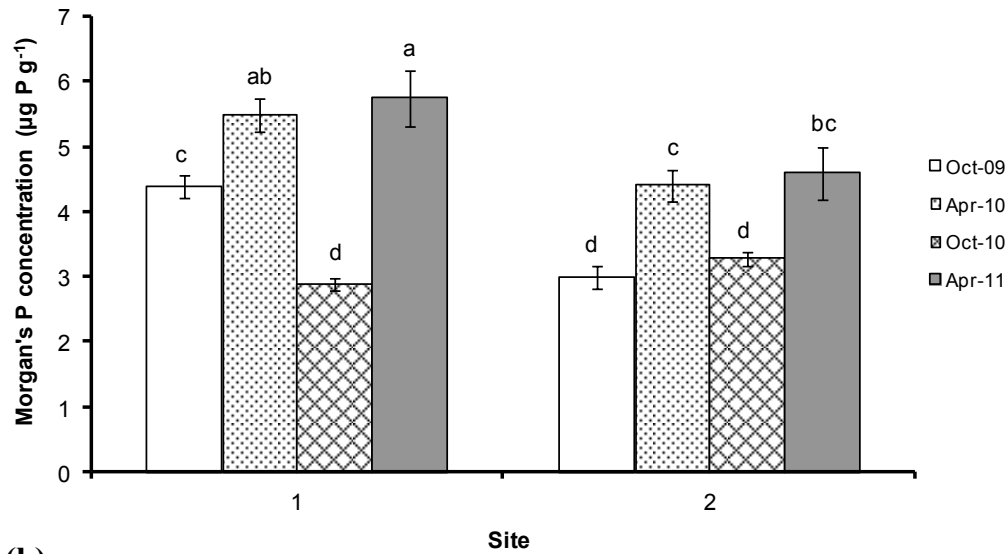
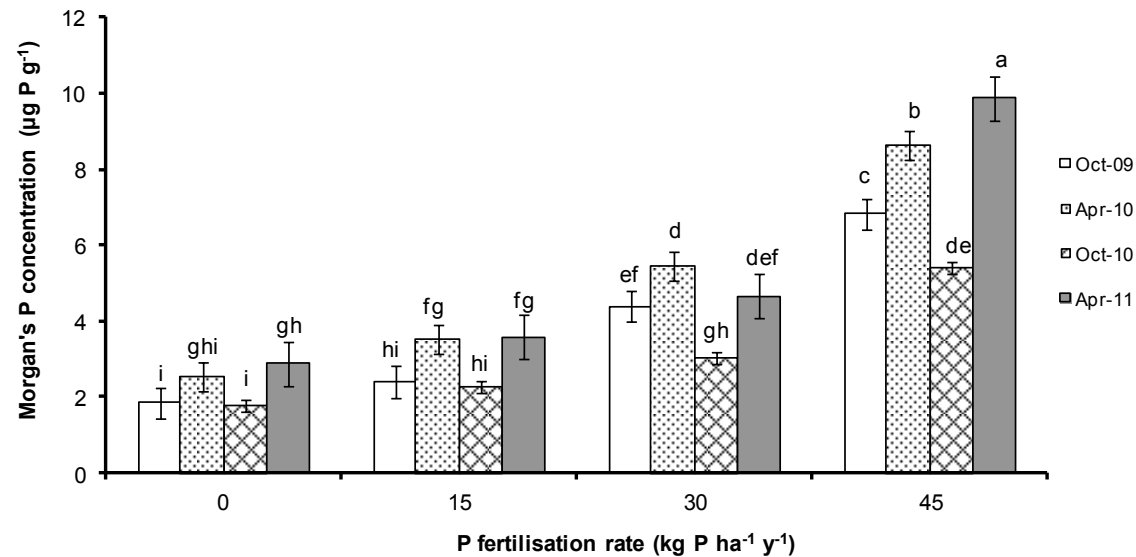


Figure 2

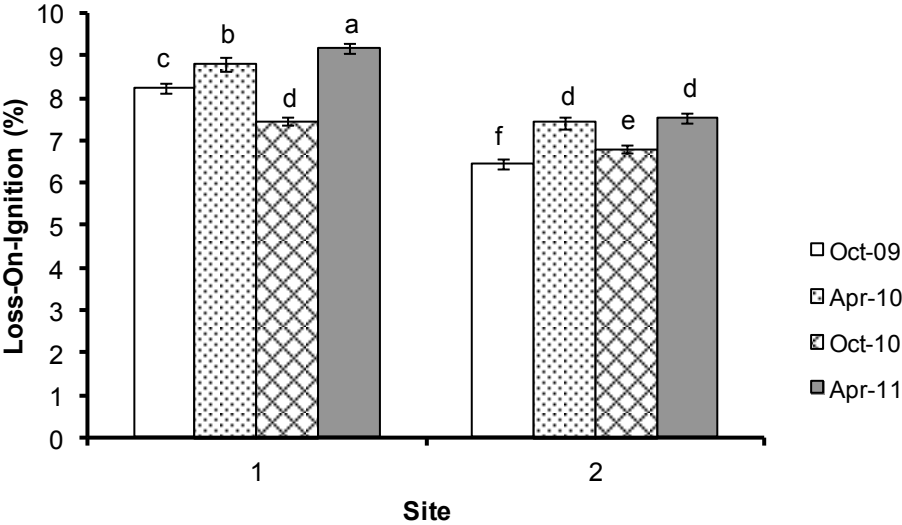
(a)



(b)

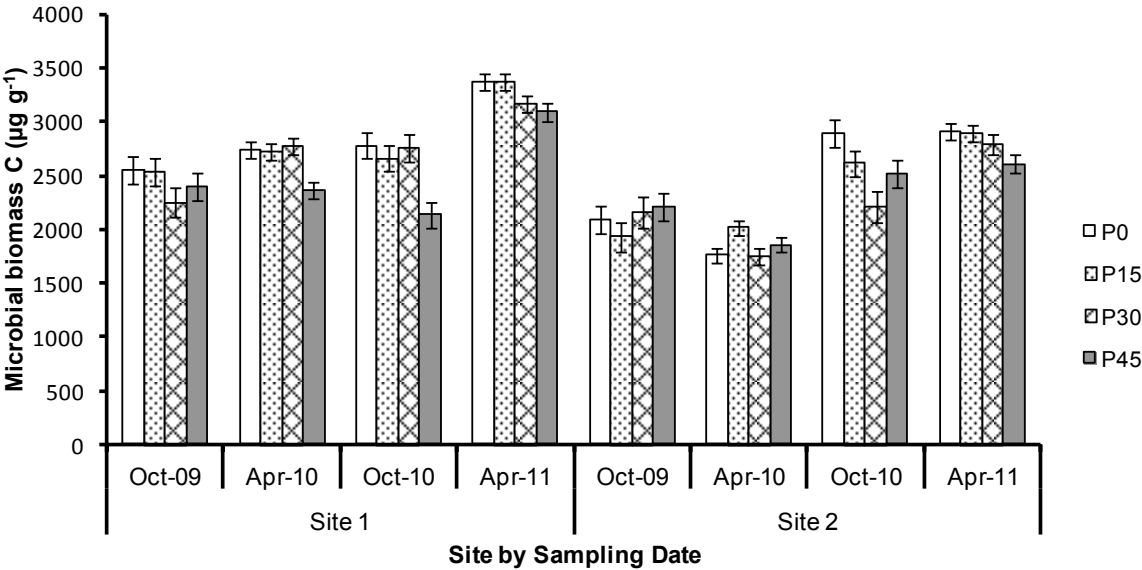


744 Figure 3



745

746 Figure 4



747

